Applicant:Adolfo J. de BoldSerial No.:10/712,335Filed:November 13, 2003Page:9 of 11

#### REMARKS

Applicant submits, through EFS-Web, a Sequence Listing in the required .txt file format and an amendment to insert the Sequence Listing and missing sequence identifiers into the specification. The sequence listing was prepared from sequences contained in the specification as filed.

I hereby state that the content of the sequence listing, submitted in accordance with 37 C.F.R. §§1.821(c) and (e), respectively, is disclosed in the originally-filed application, and as such, does not go beyond the original disclosure.

Upon entry of the present amendment, claims 1-23 will be pending. Claim 15 has been amended to insert a sequence identifier. The specification has been amended to insert sequence identifiers and correct a spelling informality. Applicant submits that no new matter has been added.

In the Office communication of March 22, 2007 (a copy of which is enclosed), the Examiner stated, "[t]he claims and instant specification lists amino acid sequences that have not been identified by [a] sequence identifier," and "Applicants are required to review the instant specification as well as the claims to comply with sequence rules" (Office communication at paragraph 2). The claims the Examiner points to with particularity (as listing amino acid sequences that have not been identified by a sequence identifier) are claims 15, 16, 18, and 19. Claim 16, for example, reads, "[t]he method of claim 14 wherein the mature BNP comprises BNP<sub>77-108</sub>."

Applicant disagrees that any sequences and sequence identifiers need to be inserted. The sequences recited in the claims were known in the art at the time the present application was filed and are not, for that reason, reproduced in the present application. One sequence (that of  $BNP_{52-76}$ ) was included in the specification (at page 18, line 28) and Applicants have now, by way of the present sequence listing and amendment, assigned a sequence identifier to that sequence and inserted the sequence identifier (SEQ ID NO:1) into the specification and claim 15.

Applicant: Adolfo J. de BoldSerial No.: 10/712,335Filed: November 13, 2003Page: 10 of 11

The present claims cover diagnostic methods and various proteins and peptides discussed in the application and recited in the claims are clearly recited by name. Applicant contends this is sufficient, as the proteins and peptides recited were known in the art at the time of filing the present application.

More specifically, the amino acid sequences discussed in the application, specifically the amino acid sequences of Homo sapiens natriuretic peptide precursor B(NPPB)(prepro BNP) and Homo sapiens natriuretic peptide precursor A(NPPA)(prepro NAF) were known in the art prior to the filing date of the instant application. Thus, the sequences of the molecules and the fragments discussed in the application would be clear and unambiguous to those skilled in the art. Sequence information for Homo sapiens natriuretic peptide precursor B(NPPB)(prepro BNP) was available from the publicly accessible National Center for Biotechnology Information (NCBI) under accession number NM 002521. Similarly, Homo sapiens natriuretic peptide precursor A(NPPA)(prepro NAF) was available under accession number NM 006172. Furthermore, with respect to the Homo sapiens natriuretic peptide precursor B(NPPB)(prepro BNP), the Examiner's attention is kindly directed to, e.g., the article by Sudoh, T. et al., entitled "Cloning and Sequence Analysis of cDNA Encoding a Precursor for Human Brain Natriuretic Peptide" published in Biochemical and Biophysical Research Communications 159 (3), 1427-1434 (1989). With respect to the *Homo sapiens* natriuretic peptide precursor A(NPPA)(prepro NAF), the Examiner is directed, e.g., to the Yang-Feng et al., reference entitled "The Pronatriodilatin Gene is Located on the Distal Short Arm of Human Chromosome 1 and on Mouse Chromosome 4" and published in the American Journal of Human Genetics 37 (6), 1117-1128 (1985). Both of these references were available to the public prior to the filing date of the instant application. Applicant submits herewith *Exhibits A* and *B*, listing the relevant NCBI accession numbers and the references associated with natriuretic peptide sequences.

Finally, Applicant requests that no charge be applied for the present Extension of Time. The Examiner states that "[s]ince the response [Applicant's last response] appears to be <u>bona</u> <u>fide</u>, but through an apparent oversight or inadvertence failed to provide a complete response;" Applicant is given one month to comply with the sequence rule. In Applicant's view, it is the Applicant:Adolfo J. de BoldSerial No.:10/712,335Filed:November 13, 2003Page:11 of 11

Office Communication that is improper. Applicant was never asked to comply with the rules regarding sequence listings. Moreover, the last Office action was a restriction requirement, to which the Applicant fully and completely replied. A Request for Refund is being submitted with the present Amendment.

A Petition for a Five-Month Extension of Time is enclosed. Please apply the required fee and any other charges or credits (as requested above and by the Request filed herewith) to deposit account 06-1050, referencing Attorney's Docket Number 14703-002001.

Respectfully submitted,

Date

-Lee Crews, Ph.D. Reg. No. 43,567

Fish & Richardson P.C. 225 Franklin Street Boston, MA 02110 Telephone: (617) 542-5070 Facsimile: (617) 542-8906

21740308.doc

	ed States Patent an	nd Trademark Office	UNITED STATES DEPAR United States Patent and Address: COMMISSIONER F P.O. Box 1450 Alexandria, Virginia 22: www.uspha.gov	Trademark Office OR PATENTS
APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/712,335	11/13/2003	Adolfo J. De Bold	14703-002001	1171
26161 7590 03/22/2007			EXAMINER	
FISH & RICHA P.O. BOX 1022			FORD, VA	NESSA L
MINNEAPOLI	S, MN 55440-1022		ART UNIT	PAPER NUMBER
			1645	
SHORTENED STATUTORY PERIOD OF RESPONSE		MAIL DATE	DELIVER	Y MODE
30 DAYS		03/22/2007	PAI	'ER

. . . . . . . . . . . . .

....

·----

... ... . .

## Please find below and/or attached an Office communication concerning this application or proceeding.

If NO period for reply is specified above, the maximum statutory period will apply and will expire 6 MONTHS from the mailing date of this communication.

-----

UNITED STATES PATENT AND TRADEMARK OFFICE



Commissioner for Patents United States Patent and Trademark Office Washington, DC 20231 www.uspid.gov

APPLICATION NO./CONTROL NO.	FILING DATE	FIRST NAMED INVENTOR /PATENT IN REEXAMINATION De Bold		ATTORNEY DOCKET NO. 14703-002001	
10/712,335					
			EXAMINER Vanessa L. Ford		
		·			
			ART UNIT	PAPER	
			1645		
			DATE MAILED		

# Please find below and/or attached an Office communication concerning this application or proceeding.

#### **Commissioner of Patents**

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 C.F.R. § 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 C.F.R. §§ 1.821-1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures. Applicant must comply with the requirements of the sequence rules (37 CFR 1.821 - 1.825) before the application can be examined under 35 U.S.C. §§ 131 and 132.

The claims and instant specification lists amino acid sequences that have not been identified by sequence identifier, e.g. SEQ ID NOs. See for example claims 15, 16, 18 and 19. See specification page 9 or 18. Applicant is required to review the instant specification as well as the claims to comply with sequence rules.

Since the response appears to be <u>bona fide</u>, but through an apparent oversight or inadvertence failed to provide a complete response, APPLICANT IS GIVEN ONE MONTH FROM THE DATE OF THIS LETTER WITHIN WHICH TO COMPLY WITH THE SEQUENCE RULES, 37 C.F.R. §§ 1.821-1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 C.F.R. § 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 C.F.R. § 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Direct the response to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Vanessa L. Ford whose telephone number is (571) 272 - 0857.

un infield PRIMARY EXAMINER

## NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
- 2. This application does not contain, as a separate part of the disclosure on paper copy, a Sequence Listing as required by 37 C.F.R. 1.821(c).
- X 3. A copy of the Sequence Listing in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
  - 4. A copy of the Sequence Listing in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up Raw Sequence Listing.
  - 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
    - 6. The paper copy of the Sequence Listing is not the same as the computer readable from of the Sequence Listing as required by 37 C.F.R. 1.821(e).
- $\overline{X}$  7. Other: No sequence Identifiers for sequences in the claims and in the specification.

### **Applicant Must Provide:**

. .... .

Ļ

- An initial or substitute computer readable form (CRF) copy of the Sequence Listing..
- An <u>initial</u> or substitute paper copy of the Sequence Listing, as well as an amendment directing its entry into the specification.

A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

PatentIn Software Program Support

. -

### PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY

## NCBI Sequence Viewer v2.0

PubMe		****	a di di conservatione di conservatione di conservatione	୬ବନ ବାହିତ୍ୟରେ କେନ୍ଦ୍ର ତେ କେ <b>ଥିକ ପ୍ରତି କେନ୍ଦ୍ର</b> ବାହିତ୍ୟରେ କେନ୍ଦ୍ର Structure	B. Prot	Taxonomy	OMIM	My NCBI [n] [Register] Books
Search	Protein	for				Co Clèar		
	Limits		Preview/Index	History	Clipboard	Detai	ils	
Display	GenPept(Full)	Show 5	Send to					
Range:	from 27	to 134	Sh	ow.whole/sequenc	e Featur	res: 🗍 SNP	🗹 CDD 📓	Refresh

□ 1: <u>NP\_002512</u>. Reports natriuretic pepti...[gi:4505433]

BLink, Conserved Domains, Links

Page 1 of 3

#### Comment Features Sequence

LOCUS DEFINITION ACCESSION VERSION DBSOURCE KEYWORDS SOURCE ORGANISM	<pre>NP_002512 108 aa linear PRI 03-JUN-2007 natriuretic peptide precursor B preproprotein [Homo sapiens]. NP_002512 REGION: 27.134 NP_002512.1 GI:4505433 REFSEQ: accession <u>NM 002521.2</u> Homo sapiens (human) <u>Homo sapiens</u> Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini; Catarrhini; Hominidae; Homo.</pre>					
REFERENCE	1 (residues 1 to 108)					
AUTHORS	Schultz, M., Kistorp, C., Langdahl, B., Raymond, I., Hildebrandt, P. and					
	Faber, J.					
TITLE	N-terminal-pro-B-type natriuretic peptide in acute hyperthyroidism					
JOURNAL	Thyroid 17 (3), 237-241 (2007)					
PUBMED	17381357					
REMARK	GeneRIF: NT-proBNP is influenced by thyroid function among healthy					
	women, as demonstrated by an inverse association between TSH and					
	NT-proBNP					
REFERENCE	2 (residues 1 to 108)					
AUTHORS	Dieplinger, B., Poelz, W., Haltmayer, M. and Mueller, T.					
TITLE	Association of adiponectin and amino terminal proBNP in peripheral					
TOTTONS	arterial disease					
JOURNAL PUBMED	Clin. Chim. Acta 377 (1-2), 192-197 (2007)					
REMARK	<u>17112494</u> GeneRIF: adiponectin and NT-proBNP were related to disease					
	severity, indicating a possible role for assessment of future morbidity and mortality in patients with peripheral arterial					
	disease					
REFERENCE	3 (residues 1 to 108)					
AUTHORS	Taskapan, M.C., Taskapan, H., Ulutas, O., Orhan, M. and Sahin, I.					
TITLE	Relationships between brain natriuretic peptide, troponin I and QT					
	dispersion in asymptomatic dialysis patients					
JOURNAL	Ren Fail 29 (2), 221-225 (2007)					
PUBMED	<u>17365940</u>					
REMARK	GeneRIF: BNP blood levels is not an indicator of heart function in					
	hemodialysis and continuous ambulatory peritoneal dialysis					
	patieents.					
REFERENCE	4 (residues 1 to 108)					
AUTHORS	Kaditis, A.G., Alexopoulos, E.I., Hatzi, F., Kostadima, E., Kiaffas, M.,					
ידי דווזי דוו	Zakynthinos, E. and Gourgoulianis, K.					
TITLE	Overnight change in brain natriuretic peptide levels in children with sleep-disordered breathing					
JOURNAL	Chest 130 (5), 1377-1384 (2006)					
PUBMED	17099013					

http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?val=4505433&from=27&to=134&view=gpwithparts

..

#### NCBI Sequence Viewer v2.0

REMARK GeneRIF: In children with snoring, overnight increase in BNP levels is correlated with severity of disturbance in respiration during sleep, which may indicate presence of nocturnal cardiac strain REFERENCE 5 (residues 1 to 108) AUTHORS Hahn, A., Schmidt, D., Hagel, K.J., Neubauer, B.A. and Katz, N. TITLE Monitoring cardiac function by B-type natriuretic peptide (BNP) in patients with infantile Pompe's disease treated with recombinant alpha-glucosidase JOURNAL Clin. Lab. 52 (11-12), 615-619 (2006) PUBMED 17175893 REMARK GeneRIF: BNP may be a valuable parameter for surveillance of cardiac function in Pompe's disease. REFERENCE 6 (residues 1 to 108) AUTHORS Suga, S., Nakao, K., Hosoda, K., Mukoyama, M., Ogawa, Y., Shirakami, G., Arai, H., Saito, Y., Kambayashi, Y., Inouye, K. et al. TITLE Receptor selectivity of natriuretic peptide family, atrial natriuretic peptide, brain natriuretic peptide, and C-type natriuretic peptide JOURNAL Endocrinology 130 (1), 229-239 (1992) PUBMED 1309330 REFERENCE 7 (residues 1 to 108) AUTHORS Hino, J., Tateyama, H., Minamino, N., Kangawa, K. and Matsuo, H. Isolation and identification of human brain natriuretic peptides in TITLE cardiac atrium JOURNAL Biochem. Biophys. Res. Commun. 167 (2), 693-700 (1990) PUBMED 2138890 REFERENCE 8 (residues 1 to 108) AUTHORS Kambayashi,Y., Nakao,K., Mukoyama,M., Saito,Y., Ogawa,Y., Shiono, S., Inouye, K., Yoshida, N. and Imura, H. TITLE Isolation and sequence determination of human brain natriuretic peptide in human atrium FEBS Lett. 259 (2), 341-345 (1990) JOURNAL 2136732 PUBMED REFERENCE 9 (residues 1 to 108) AUTHORS Seilhamer, J.J., Arfsten, A., Miller, J.A., Lundquist, P., Scarborough, R.M., Lewicki, J.A. and Porter, J.G. TITLE Human and canine gene homologs of porcine brain natriuretic peptide JOURNAL Biochem. Biophys. Res. Commun. 165 (2), 650-658 (1989) PUBMED 2597152 REFERENCE 10 (residues 1 to 108) AUTHORS Sudoh, T., Maekawa, K., Kojima, M., Minamino, N., Kangawa, K. and Matsuo.H. TITLE Cloning and sequence analysis of cDNA encoding a precursor for human brain natriuretic peptide JOURNAL Biochem. Biophys. Res. Commun. 159 (3), 1427-1434 (1989) PUBMED 2522777 COMMENT REVIEWED REFSEQ: This record has been curated by NCBI staff. The reference sequence was derived from AJ708502.1, M25296.1 and BC025785.1. Summary: This gene is a member of the natriuretic peptide family and encodes a secreted protein which functions as a cardiac hormone. The protein undergoes two cleavage events, one within the cell and a second after secretion into the blood. The protein's biological actions include natriuresis, diuresis, vasorelaxation, inhibition of remin and aldosterone secretion, and a key role in cardiovascular homeostasis. A high concentration of this protein in the bloodstream is indicative of heart failure. Mutations in this gene have been associated with postmenopausal osteoporosis. Publication Note: This RefSeq record includes a subset of the

publications that are available for this gene. Please see the Entrez Gene record to access additional publications.

http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?val=4505433&from=27&to=134&view=gpwithparts

11/06/2007

NCBI Sequence Viewer	v2.0
FEATURES	Location/Qualifiers
source	1108
	/organism="Homo sapiens"
	/db xref="taxon:9606"
	/chromosome="1"
	/map="1p36.2"
Protein	<1108
1.00011	/product="natriuretic peptide precursor B preproprotein"
	/note="brain type natriuretic peptide; natriuretic
	protein"
proprotein	1108
proprocern	
	/product="natriuretic peptide precursor B proprotein"
	/note="proBNP; gamma-brain natriuretic peptide"
Do and and	/calculated_mol_wt=11906
Region	20102
	/region_name="ANP"
	/note="Atrial natriuretic peptide; pfam00212"
	/db_xref="CDD: <u>64095</u> "
mat_peptide	77108
	/product="natriuretic peptide precursor B"
	/note="brain natriuretic peptide 32"
	/calculated_mol_wt=3466
CDS	<1108
	/gene="NPPB"
	/coded_by="NM_002521.2:103507"
	<pre>/GO_component="extracellular region [pmid 14960748];</pre>
	<u>extracellular space</u> [pmid <u>7601467</u> ] "
	/GO_function="diuretic hormone activity [pmid 14960748]"
	<pre>/GO_process="blood pressure regulation [pmid 12727915];</pre>
	cell surface receptor linked signal transduction [pmid
	12727915]; diuresis [pmid 14737067]; fluid secretion [pmid
	14960748]; natriuresis [pmid 14960748]; negative
	regulation of anglogenesis [pmid 14737067]; negative
	regulation of cell growth [pmid 12727915]; regulation of
	blood vessel size; regulation of vascular permeability
	[pmid 14737067]; regulation of vasodilation [pmid
	7601467] "
	/db_xref="CCDS:CCDS140.1"
	/db_xref="GeneID:4879"
	/db xref="HGNC:7940"
	/db_xref="HPRD:08974"
	/db xref="MIM:600295"
ORIGIN	
1 hplgspgsas	dletsglqeq rnhlqgklse lqveqtslep lqesprptgv wksrevateg
61 irghrkmvly	tlraprspkm vqgsgcfgrk mdrissssgl gckvlrrh
//	

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

Apr 17 2007 11:10:07

• •

.

Page 3 of 3

NCBI Sequen	nce Viewer v2.0	Page 1 of 4
PubMed	CCCCA CLASSAT SAGACTICGASCOCTAGLOGATCGCGATCCCCGCCCC COLATIATATAGCTCGATCATCATCA ILCTCT TAT CACCOC SCATGOR STATATACACACAC CCCGCCGCGATAGCATCACTGATCTA BLACCCA CLASSAT SAGACTACTICGCATACGIGTECATEGCCGCGCGCGCGCGCCTA CACAGACTACTACTACTACTACCAATTCGGCGCGCGCGCG	My NCBI [Sign In] [Register] OMIM Books
Sameh Cor		Summer Boons
Search Cor	eNucleotide for Go	
	Limits Preview/Index History Clipboard Details	
Display Gen	Bank 🔣 Show 5 💽 Send to 🔀 Hide: 🗆 sequence 🗔 all but gene, CDS and mF	INA features
Range: from	begin to end I Reverse complemented strand Features: SNP I STS	Refresh
□1: <u>NM_0</u> (	06172. Reports Homo sapiens natr[gi:141803366]	Links
<u>Comment</u>	Features Sequence	
LOCUS DEFINITION ACCESSION	NM_006172 840 bp mRNA linear PRI 03-JUN-2007 Homo sapiens natriuretic peptide precursor A (NPPA), mRNA. NM_006172	
VERSION KEYWORDS	NM_006172.2 GI:141803366	
SOURCE	Homo sapiens (human)	
ORGANISM	<u>Homo sapiens</u> Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini;	
REFERENCE	Catarrhini; Hominidae; Homo. 1 (bases 1 to 840)	
AUTHORS	Batlle, M., Roig, E., Perez-Villa, F., Lario, S., Cejudo-Martin, P.,	
	Garcia-Pras, E., Ortiz, J., Roque, M., Orus, J., Rigol, M., Heras, M.,	
	Ramirez, J. and Jimenez, W.	
TITLE	Increased expression of the renin-angiotensin system and mast cell density but not of angiotensin-converting enzyme II in late stages	
JOURNAL	of human heart failure J. Heart Lung Transplant. 25 (9), 1117–1125 (2006)	
PUBMED	16962475	
REMARK	GeneRIF: Both ANP and BNP expression were higher in heart failure than in control samples.	
REFERENCE	2 (bases 1 to 840)	
AUTHORS TITLE	He,X.L., Dukkipati,A. and Garcia,K.C. Structural determinants of natriuretic peptide receptor specificity and degeneracy	
JOURNAL	J. Mol. Biol. 361 (4), 698-714 (2006)	
PUBMED	16870210	
REFERENCE	3 (bases 1 to 840)	
AUTHORS	Rubattu,S., Bigatti,G., Evangelista,A., Lanzani,C., Stanzione,R., Zagato,L., Manunta,P., Marchitti,S., Venturelli,V., Bianchi,G., Volpe,M. and Stella,P.	
TITLE	Association of atrial natriuretic peptide and type a natriuretic peptide receptor gene polymorphisms with left ventricular mass in	
TOTIONIAT	human essential hypertension	
JOURNAL PUBMED	J. Am. Coll. Cardiol. 48 (3), 499-505 (2006) 16875975	
REMARK	GeneRIF: ANP/NPRA system significantly contributes to ventricular	
	remodeling in human essential hypertension.	
REFERENCE	4 (bases 1 to 840)	
AUTHORS	Lim,J., Hao,T., Shaw,C., Patel,A.J., Szabo,G., Rual,J.F., Fisk,C.J., Li,N., Smolyar,A., Hill,D.E., Barabasi,A.L., Vidal,M.	
TITLE	and Zoghbi,H.Y. A protein-protein interaction network for human inherited ataxias and disorders of Purkinje cell degeneration	
JOURNAL	Cell 125 (4), 801-814 (2006)	
PUBMED	<u>16713569</u>	
REFERENCE	5 (bases 1 to 840)	

http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=nuccore&id=141803366

.

#### NCBI Sequence Viewer v2.0

AUTHORS Dedoussis, G.V., Maumus, S., Skoumas, J., Choumerianou, D.M., Pitsavos, C., Stefanadis, C. and Visvikis-Siest, S. TITLE Natriuretic peptide Val7Met substitution and risk of coronary artery disease in Greek patients with familial hypercholesterolemia JOURNAL J. Clin. Lab. Anal. 20 (3), 98-104 (2006) PUBMED 16721833 REMARK GeneRIF: The 664A allele of the Atrial Natriuretic Peptide (ANP) polymorphism is associated with lower levels of ApoA1 and HDL-C in Familial Hypercholesterolemia patients, but not with Coronary Artery Disease risk. REFERENCE (bases 1 to 840) 6 AUTHORS Suga, S., Nakao, K., Hosoda, K., Mukoyama, M., Ogawa, Y., Shirakami, G., Arai, H., Saito, Y., Kambayashi, Y., Inouye, K. et al. TITLE Receptor selectivity of natriuretic peptide family, atrial natriuretic peptide, brain natriuretic peptide, and C-type natriuretic peptide JOURNAL Endocrinology 130 (1), 229-239 (1992) PUBMED 1309330 REFERENCE (bases 1 to 840) 7 AUTHORS Vanneste, Y., Michel, A. and Deschodt-Lanckman, M. Hydrolysis of intact and Cys-Phe-cleaved human atrial natriuretic TITLE peptide in vitro by human tissue kallikrein JOURNAL Eur. J. Biochem. 196 (2), 281-286 (1991) 1826098 PUBMED REFERENCE (bases 1 to 840) 8 AUTHORS Porter, J.G., Arfsten, A., Fuller, F., Miller, J.A., Gregory, L.C. and Lewicki, J.A. TITLE Isolation and functional expression of the human atrial natriuretic peptide clearance receptor cDNA JOURNAL Biochem. Biophys. Res. Commun. 171 (2), 796-803 (1990) PUBMED 2169733 REFERENCE 9 (bases 1 to 840) AUTHORS Yandle, T.G., Brennan, S.O., Espiner, E.A., Nicholls, M.G. and Richards, A.M. TITLE Endopeptidase-24.11 in human plasma degrades atrial natriuretic factor (ANF) to ANF(99-105/106-126) JOURNAL Peptides 10 (4), 891-894 (1989) PUBMED 2531377 REFERENCE 10 (bases 1 to 840) AUTHORS Yang-Feng, T.L., Floyd-Smith, G., Nemer, M., Drouin, J. and Francke, U. TITLE The pronatriodilatin gene is located on the distal short arm of human chromosome 1 and on mouse chromosome 4 JOURNAL Am. J. Hum. Genet. 37 (6), 1117-1128 (1985) PUBMED 2934979 PROVISIONAL REFSEQ: This record has not yet been subject to final COMMENT NCBI review. The reference sequence was derived from M30262.1. On Apr 5, 2007 this sequence version replaced gi:23510318. Sequence Note: removed 5 bases from the 5' end that did not align to the reference genome assembly. Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications. FEATURES Location/Qualifiers source 1..840 /organism="Homo sapiens" /mol\_type="mRNA" /db xref="taxon:9606" /chromosome="1" /map="1p36.21" gene 1..840 /gene="NPPA"

http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=nuccore&id=141803366

			/note="nat:	riuretic per	otide precu	csor A; sync	onyms: ANF,		
			ANP, PND, CDD-ANF"						
			/db xref="GeneID:4878"						
			/db xref="HGNC:7939"						
			/db xref="HPRD:00164"						
			/db xref="MIM: 108780"						
	CDS		95550						
			/gene="NPPA"						
			/GO_component="extracellular region [pmid 14718574]"						
			/GO function="hormone_activity"						
			/GO process="blood pressure regulation [pmid 6230082]						
			[pmid 6547996]; physiological process; regulation of blood						
			vessel size		Judgidur pr	<u></u> , <u></u>			
					In• atrial 1	atriuretic	peptide"		
			<pre>/note="pronatriodilatin; atrial natriuretic peptide" /codon start=1</pre>						
			/product="natriuretic peptide precursor A"						
			/protein_id=" <u>NP 006163.1</u> "						
				JI:23510319					
				CCDS: CCDS139					
				GeneID:4878					
			/db xref="l						
			. —	HPRD:00164"					
				MIM: 108780"			·		
					N70 PT.T.T.T. 7 POI				
			/translation="MSSFSTTTVSFLLLLAFQLLGQTRANPMYNAVSNADLMDFKNLL						
			DHLEEKMPLEDEVVPPQVLSEPNEEAGAALSPLPEVPPWTGEVSPAQRDGGALGRGPW						
	sian	peptide	DSSDRSALLKSKLRALLTAPRSLRRSSCFGGRMDRIGAQSGLGCNSFRY" 95169						
	<u></u>	Joperue	/gene="NPPA"						
			/gene="NPPA" /note="cardiodilatin-atrial natriuretic factor signal						
			peptide"						
	mat 7	peptide	170370						
	mac j	Jeperde	/gene="NPPA"						
	mat r	peptide	/product="cardiodilatin" 464547						
	mac j	Septide	/gene="NPPA"						
			/product="atrial natriuretic factor"						
	omo		349599						
	STS								
			/gene="NPPA"						
			/standard_name="GDB:226664"						
	STS		/db_xref="UniSTS: <u>156242</u> "						
	515								
			/gene="NPPA"						
			/standard_name="NPPA"						
	ama		/db_xref="UniSTS: <u>253991</u> "						
	STS								
			/gene="NPPA						
			/standard_name="SHGC-74238" /db xref="UniSTS:43068"						
ODTOT	'NT		/db_xrei="(	mists: 43068	<u>s</u> "				
ORIGI			atogaaaaa	6 <i>a</i> 6 <i>a</i>					
		agggacagac							
	101	agcaagcagt	yyactgetee	Lugacgacge	cagcatgage	LCCTTCTCCA	ccaccaccgt		
	101	gagetteete	tampart	cattecaget	cetaggteag	accagageta	atcccatgta		
		caatgccgtg							
		aaagatgcct							
		ageggggget							
		agcccagaga							
		cctcctaaaa							
		ctgcttcggg							
		ccggtactga							
		agaggeteet							
		tgatcatccc							
		agataaagtg							
	10T	gtagaacctc	acceguiade	yaaayuyytt	uyaaayuyaa	Laaacticag	Laucalyyac		

### Disclaimer | Write to the Help Desk NCBI | NLM | NIH

.

Apr 17 2007 11:10:07

٠.

### 11/06/2007